To get ce en roc ce ase ce mo cer en THE ATC CCG GCC CCT GCT CCA GGC CTC ACT GTG CAA GTG CTG CTG TC 29 L L L L M P V N P Q R L P R M Q 97 CTO CTO CTT CTO ATO CCT GTC CAT CCC CAG AGG TTG CCC CGG ATO CAG GAG GAT TCC CCC TTG GGA GGA GGA GGC TET TCT GGG GAA GAT GAC CCA CTG O E E D L P B E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S P R E D S F P G E E D L P G E E D L P G E CCA CCC OGA GAG GAT CTA CCT OGA GAG GAT CTA CCT OGA GAG GAG GAT CTA CCT GAA GTT AAG CCT AAA TCA GAA GAA GAG GGC TCC CTG AND THA GAS ONT CTA CCT ACT OTT GAG GGL CCL GGA GAL CCT CAR GRA CCC CAG AAT AAT OCC CAC AGG GAC AAA GAA GGG GAT GAC CAG AGT CAT TOG COC TAT GOA GOC GAC CCG CCC TOG CCC CGG GTG TCC CCA GCC TGC A O R P O S P V D I R P O L A A OCCU COC COC CAG CTC CCC CAG CTC CCC F C P A L R P L E L L G F G L P TTC TGC CCG CCC CTG GAA CTC CTG GGC TTC CAG CTC CCG P L P E L R L R M N G H B V Q L CCG CTC CCA GAA CTG CCC CAAC AAT GGC CAC AAT GTG CAA CTG T L P P G L E M A L G P G R E Y ACC CTG CCT CCT GGG CTG GGG CT R A L Q L H L H W 6 A A G R P G CGG GCT CTG CAG CTG CAG GCG GCT GCA GGT CGG GGG S E H T V E G H R F P A E I M V TCG GAG CAC ACT GTG GAA GGC CAC CGT TTC CCT GCC GAG ATC CAC GTG OTT CAC CTC AGC ACC GCC TIT GCC AGA GTT GAC GAG GCC TTG GGG GCC CCG GGA GGC CTG GCC GTG TTG GCC GCC TTT CTG GAG GAG GGC CCG GAA GAA AAC AGT OCC TAT GAG CAG TTG CTC TCT CGC TTG GAA GAA ATC CCT GAG GAA OGC TCA GAG ACT CAG GTC CCA GGA CTG GAC ATA TCT GGA CTC CTO CCC TCT GAC TTC AGC COC TAC TTC CAA TAT GAG GOO TCT CTG ACT 332 ACA CCG CCC TOT OCC CAG GOT OTC ATC TOG ACT OTG TIT AAC CAG ACA 348 OTG ATG CTG AGT GGT AAG CAG CTC CAC ACC CTC TCT GAC ACC CTG TGG 364 GA CCT GGT GAC TUT CGG CTA CAG CTG AAC TTC CGA GCG ACG CAG CCT THE AAT GOO COA GTO ATT GAG GCC TCC TTC CCT GGA GTO GAA AND ACT CCT COO OCT OCT OAG CCA OTC CAO CTG AAT TCC TGC CTG OCT OCT 412 OF OAC ATC CTA OCC CTG GTT TTT OGC CTC CTT TTT OCT GTC ACC AGC 428 1296 1297 OTC GCG TTC CTT GTG CAG ATG AGA AGG CAG CAG AGA AGG GGA AGC AAA 1344 445 GO GOT GTO AGC TAC COC CCA GCA GAG GTA GCC GAG ACT GGA GCC TAG 460 1392 AGG CTG GAT CIT GGA GAA TOT GAG AAG CCA GCC AAA GGC ATC TGA GGG GGA GCC GGT AAC TOT CCT GTC CTG CTC ATT ATG CCA CTT CCT TTT AAC TGC CAA GAA ATT TIT TAA AAT AAA TAT TAT TAA T 1440 1468 1522

92 286

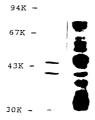
108 336

124

268 816

FIG. 1

## А В



20.1K -

FIG. 2.

# A B C D



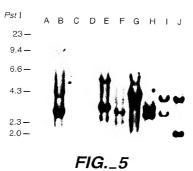
FIG. 3.

### ABCDEF

1.5 kb -



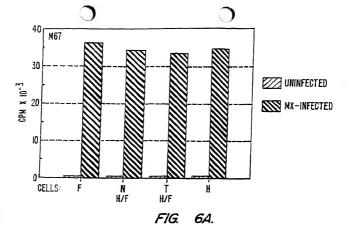
FIG. 4.

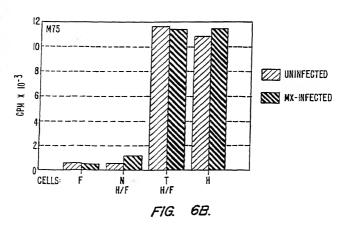


+

A B C D E F G H E J

FIG. 5





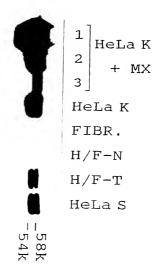


FIG. 7.



FIG. 8

# F G H I ABCDE

K L M N O P

58k-54k<sup>-</sup>



F16. 9.

+ME OME

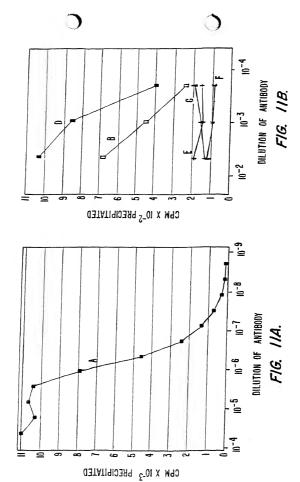
A B A B

153k-



58k-54k- **8**- **9** 

FIG. 10.



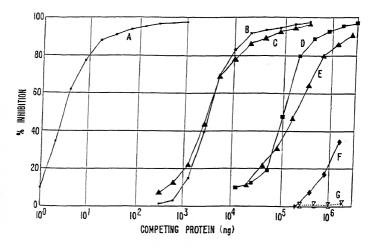


FIG. 12.

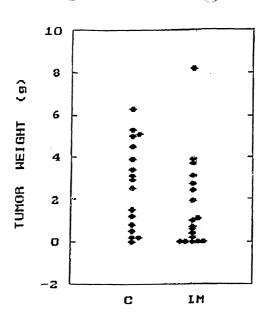


FIG. 14

1 ggatectgtt gac-egtgac ettacececa accetgtge etetgaaaca tgagetgtgt 61 ccactcaggg ttaaatggat taagggcggt gcaagatgtg ctttgttaaa cagatgcttg 121 aaggeageat getegttaag agteateace aateeetaat eteaagtaat eagggacaca 181 aacactgogg aaggoogcag ggtoctotgc ctaggaaaac cagagacett tgttcacttg 241 tttatctgac cttccctcca ctattgtcca tgaccctgcc aaatccccct ctgtgagaaa 361 aaaaaaaaa gacttacgaa tagttattga taaatgaata gctattggta aagccaagta 421 aatgatcata ticaaaacca gacggccatc atcacagctc aagtctacct gatttgatct 481 etttateatt gteattettt ggatteacta gattagteat cateeteaaa atteteecee 541 aagttetaat taegtteeaa acatttaggg gttacatgaa gettgaacet actacettet 601 ttgettttga gecatgagtt gtaggaatga tgagtttaca cettacatge tggggattaa 661 tttaaacttt acctctaagt cagttgggta qcctttgget tatttttgta gctaattttg 721 tagttaatgg atgcactgtg aatcttgcta tgatagtttt cctccacact ttgccactag 781 gggtaggtag gtactcagtt ttcagtaatt gcttacctaa gaccctaagc cctatttctc 841 ttgtactggc ctttatctgt aatatgggca tatttaatac aatataattt ttggagtttt 901 titigtitigti tigtitigtitig tittititigag acggagitett geatetigtea tigceeagget 961 ggagtageag tggtgccate teggeteact geaageteea cetecegagt teacgecatt 1021 tteetgeete ageeteega gtagetggga etacaggege cegecaccat geeeggetaa 1081 ttttttgtat ttttggtaga gacggggttt caccgtgtta gccagaatgg tctcgatctc 1141 ctgacttegt gatecacecg ceteggeete ecaaagttet gggattacag gtgtgageca 1201 cogcacctog ccaatttttt gagtetttta aagtaaaaat atgtettgta agetggtaac 1261 tatogtacat tteettttat taatgtggtg etgaeggtea tataggttet tttgagtttg 1321 gcatgcatat gctacttttt gcagtccttt cattacattt ttctctcttc atttgaagag 1381 catgitatat cittitagcit cactiggcit aaaaggitci cicattagcc taacacagtg 11441 teattgttgg taccaettgg atcataagtg gaaaaacagt caagaaattg cacagtaata 1501 cttgtttgta agagggatga ttcaggtgaa tctgacacta agaaactccc ctacctgagg 1561 tetgagatte etetgacatt getgtatata ggetttteet ttgacageet gtgactgegg 11621 actatttttc ttaagcaaga tatgctaaag ttttgtgagc ctttttccag agagaggtct 1681 catatetgca tcaagtgaga acatataatg tetgcatgtt tccatattte aggaatgttt 1741 gettgtgttt tatgetitta tatagacagg gaaacttgtt cetcagtgae ceaaaagagg 1801 tgggaattgt tattggatat catcattgge ceaegettte tgacettgga aacaattaag 1861 ggttcataat ctcaattctg tcagaattgg tacaagaaat agctgctatg tttcttgaca 1921 ttccacttgg taggaaataa gaatgtgaaa ctcttcagtt ggtgtgtgtc cct?gttttt 1 1981 ttgcaatttc cttcttactg tgttaaaaaa aagtatgatc ttgctctgag aggtgaggca 2041 ttcttaatca tgatctttaa agatcaataa tataatcctt tcaaggatta tgtctttatt 2101 ataataaaga taatttgtct ttaacagaat caataatata atcccttaaa ggattatatc ₩2161 tttgctgggc gcagtggctc acacctgtaa tcccagcact ttgggtggcc aaggtggaag 2221 gatcaaattt geetaettet atattatett etaaageaga atteatetet etteeeteaa 2281 tatgatgata ttgacagggt ttgccctcac tcactagatt gtgagctcct gctcagggca 2341 ggtagegttt titgttittg titttgttit tettittga gacagggtet tgetetgtca 2401 cccaggocag agtgcaatgg tacagtotca gotcaetgca goctcaaccg cotcggotca 2461 aaccatcate ecattteage etectgagta getgggacta caggeacatg ceattacace 2521 tggctaattt ttttgtatit ctagtagaga cagggtttgg ccatgttgcc cgggctggtc 2581 tegaacteet ggaeteaage aatecacea ceteageete ceaaaatgag ggaeegtgte 2641 ttattcattt ccatgtccct agtccatagc ccagtgctgg acctatggta gtactaaata 2701 aatatttgtt gaatgcaata gtaaatagca tttcagggag caagaactag attaacaaag 2761 gtggtaaaag gtttggagaa aaaaataata gtttaatttg gctagagtat gagggagagt 2821 agtaggagac aagatggaaa ggtctcttgg gcaaggtttt gaaggaagtt ggaagtcaga 2881 agtacacaat gtgcatatcg tggcaggcag tggggagcca atgaaggctt ttgagcagga 2941 gagtaatgtg ttgaaaaata aatataggtt aaacctatca gagcccctct gacacataca 3001 cttgetttte attcaagete aagtttgtet eccacatace cattacttaa etcaceeteg

3061 ggetecceta gea stgee ctacetettt acctgette lggtggagtc agggatgtat 3121 acatgagetg ctttccctct cagccagagg acatggggggg ccccagetce cctgcctttc 3181 cccttctgtg cctggagctg ggaagcaggc cagggttagc tgaggctggc tggcaagcag 3241 ctgggtggtg ccagggagag cctgcatagt gccaggtggt gccttgggtt ccaagctagt 3301 ccatggcccc gataaccttc tgcctgtgca cacacctgcc cctcactcca cccccatcct 3361 agetttggta tgggggagag ggcacagggc cagacaaacc tgtgagactt tggctccatc 3421 tetgeaaaag ggegetetgt gagteageet geteeeetee aggettgete eteeeeeaee 3481 cagetetegt ticcaatgea cotacagee otacacage tgtgetggga caceccACAG 3541 TCAGCCGCAT GGCTCCCTG TGCCCCAGCC CCTGGCTCCC TCTGTTGATC CCGGCCCCTG 3601 CTCCAGGCCT CACTGTGCAA CTGCTGCTGT CACTGCTGCT TCTGGTGCCT GTCCATCCCC 3661 AGAGGTTGCC CCGGATGCAG GAGGATTCCC CCTTGGGAGG AGGCTCTTCT GGGGAAGATG 3721 ACCCACTGGG CGAGGAGGAT CTGCCCAGTG AAGAGGATTC ACCCAGAGAG GAGGATCCAC 3781 CCGGAGAGGA GGATCTACCT GGAGAGGAGG ATCTACCTGG AGAGGAGGAT CTACCTGAAG 3881 TTAAGCCTAA ATCAGAAGAA GAGGGCTCCC TGAAGTTAGA GGATCACCT ACTGTGAGG 3901 CTCCTGGAGA TCCTCAAGAA CCCCAGAATA ATGCCCACCAG GGACAAAGAA GALBBGIEGI 3961 catcaatoto caaatocagg ttocaggagg ttoatgacto cootoccata coccagoota 4021 ggetetgtte acteagggaa ggaggggaga etgtacteec cacagaagee ettecagagg 4081 teccatacca atatececat ecceaetete ggaggtagaa agggacagat gtggagagaa 4141 aataaaaagg gtgcaaaagg agaqaggtga gctggatgag atgggagaga agggggaggc 4201 tggagaagag aaagggatga gaactgcaga tgagagaaaa aatgtgcaga cagaggaaaa 4261 aaataggtgg agaaggagag tcagagagtt tgaggggaag agaaaaggaa agcttgggag 4321 gtgaagtggg taccagagac aagcaagaag agctggtaga agtcatctca tettaggeta 4381 caatgaggaa ttgagaccta ggaagaaggg acacagcagg tagagaaacg tggcttettg 4441 actoccaago caggaatttg gggaaagggg ttggagacca tacaaggcag agggatgagt 4501 ggggagaaga aagaagggag aaaggaaaga tggtgtactc actcatttgg gactcaggac 4561 tgaagtgccc actcactttt ttttttttt tttttgagac aaactttcac ttttgttgcc 4521 caggetggag tgcaatggcg cgatetegge teactgcaac etccacetee cgggttcaag 4681 tgatteteet geeteageet etagecaagt agetgegatt acaggeatge geeaceaege 4741 coggetaatt tttgtatttt tagtagagac ggggtttcgc catgttggtc aggetggtct 4801 cgaactcetg atctcaggtg atccaaccac cetggeetec caaagtgetg ggattatagg 4861 cgtgagccac agcgcctggc ctgaagcagc cactcacttt tacagaccct aagacaatga 4921 ttgcaagetg gtaggattge tgtttggccc acccagetge ggtgttgagt ttgggtgegg 4981 tetectgtge tttgcacetg gcccgettaa ggcatttgtt accegtaatg etectgtaag 5041 gcatctgcgt ttgtgacatc gttttggtcg ccaggaaggg attggggctc taagcttgag 5101 cggttcatcc ttttcattta tacagcGGAT GACCAGAGTC ATTGGCGCTA TGGAGgtgag 5161 acacccacco gotgoacaga occaatotgg gaacccagot otgtggatot occotacago 5221 cgtccctgaa cactggtccc gggcgtccca cccgccgccc accgtcccac cccctcacct 5281 tttctacccg ggttccctaa gttcctgacc taggcgtcag acttcctcac tatactctcc 5341 caccecagGC GACCCGCCCT GGCCCCGGGT GTCCCCAGCC TGCGCGGGCC GCTTCCAGTC 5401 CCCGGTGGAT ATCCGCCCCC AGCTCGCCGC CTTCTGCCCG GCCCTGCGCC CCCTGGAACT 5461 CCTGGGCTTC CAGCTCCCGC CGCTCCCAGA ACTGCGCCTG CGCAACAATG GCCACAGTGG 5521 tgagggggtc tccccgccga gacttgggga tggggcgggg cgcagggaag ggaaccgtcg 5581 cgcagtgcct gecegggggt tgggctggcc ctacegggeg gggceggete acttgcctct 5641 ccctacgcag TGCAACTGAC CCTGCCTCCT GGGCTAGAGA TGGCTCTGGG TCCCGGGCGG 5701 GAGTACCGGG CTCTGCAGCT GCATCTGCAC TGGGGGGGCTG CAGGTCGTCC GGGCTCGGAG 5761 CACACTGTGG AAGGCCACCG TTTCCCTGCC GAGgtgageg eggaetggee gagaagggge 5821 aaaggagegg ggcggaeggg ggceagagae gtggeeetet cetaceeteg tgteetttte 5881 agATCCACGT GGTTCACCTC AGCACCGCCT TTGCCAGAGT TGACGAGGCC TTGGGGCGCC 5941 CGGGAGGCCT GGCCGTGTTG GCCGCCTTTC TGGAGGtacc agatectgga caccectac 6001 teccegettt cecateceat geteeteeeg gaetetateg tggagecaga gaeeceatee 6061 cagcaagete acteaggeee etggetgaca aacteattea egeactgitt giteattiaa 6121 cacccaetgt gaaccaggca ccagcccca acaaggatte tgaagetgta ggteettgee 6181 tetaaggage ccacagecag tgggggagge tgacatgaca gacacatagg aaggacatag

FIG. 15b

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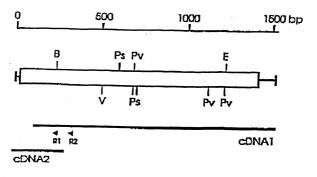


FIG. 16

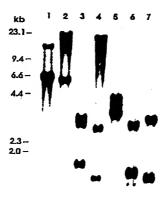


FIG. 17

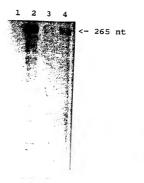


FIG 18a

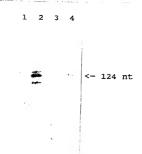


FIG. 18b

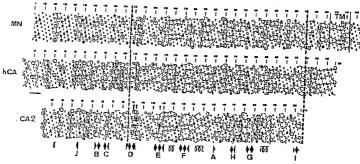


Fig. 19a

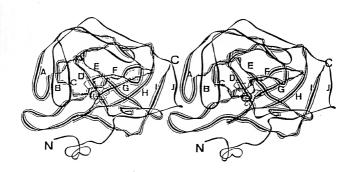
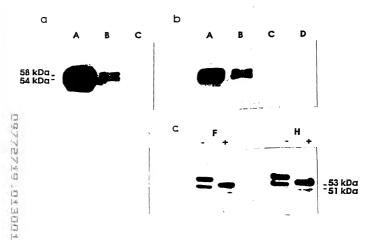


Fig. 19b

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FIG. 21

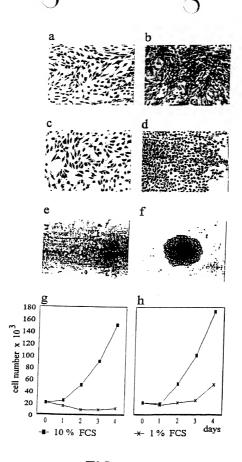


FIG. 22

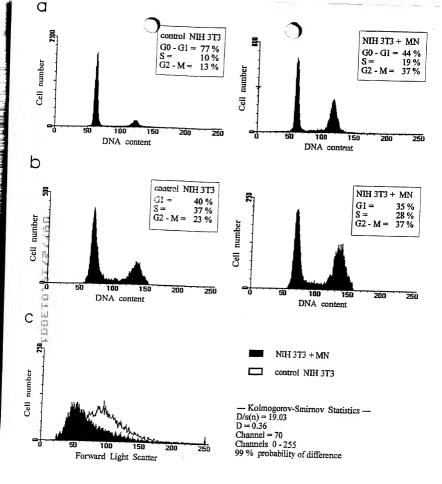


FIG. 23

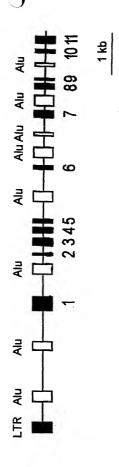
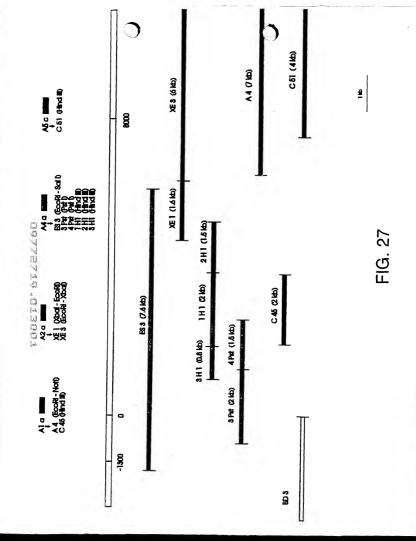


FIG. 24

| -506 | CTTGCTTTTC                        | ATTCAAGCTC                                                                                                         | AAGTTTGTCT                                                                                                                                                                                      | CCCACATACC                                                                                                                                                                                                                                                                                       | CATTACTTAA                                                                                                                                                                                                                                                                                                                                  | CTCACCCTCG                                                                                                                                                                                                                                                                                                                                                                     |
|------|-----------------------------------|--------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| -446 | GGCTCCCCTA<br>AP2                 | GCAGCCTGCC<br>AP2                                                                                                  | CTACCTCTTT                                                                                                                                                                                      | ACCTGCTTCC                                                                                                                                                                                                                                                                                       | TGGTGGAGTC                                                                                                                                                                                                                                                                                                                                  | AGGGATGTAT                                                                                                                                                                                                                                                                                                                                                                     |
| -386 | ACATGAGCTG                        | CTTTCCCTCT                                                                                                         | CAGCCAGAGG                                                                                                                                                                                      | ACATGGGGGG                                                                                                                                                                                                                                                                                       | CCCCAGCTCC                                                                                                                                                                                                                                                                                                                                  | CCTGCCTTTC                                                                                                                                                                                                                                                                                                                                                                     |
| -326 | CCCTTCTGTG                        | CCTGGAGCTG                                                                                                         | GGAAGCAGGC                                                                                                                                                                                      | CAGGGTTAGC                                                                                                                                                                                                                                                                                       | TGAGGCTGGC                                                                                                                                                                                                                                                                                                                                  | TGGCAAGCAG                                                                                                                                                                                                                                                                                                                                                                     |
| -266 | CTGGGTGGTG                        | CCAGGGAGAG                                                                                                         | CCTCCATAGT                                                                                                                                                                                      | GCCAGGTGGT                                                                                                                                                                                                                                                                                       | GCCTTGGGTT                                                                                                                                                                                                                                                                                                                                  | CCAAGCTAGT                                                                                                                                                                                                                                                                                                                                                                     |
| -206 | CCATGGCCCC                        | GATAACCTTC                                                                                                         | TGCCTGTGCA                                                                                                                                                                                      | CACACCTGCC                                                                                                                                                                                                                                                                                       | CCTCACTCCA                                                                                                                                                                                                                                                                                                                                  | CCCCCATCCT                                                                                                                                                                                                                                                                                                                                                                     |
| -146 | AGCTTTGGTA                        | TGGGGGAGAG                                                                                                         | GGCACAGGGC                                                                                                                                                                                      | CAGACAAACC                                                                                                                                                                                                                                                                                       | TGTGAGACTT                                                                                                                                                                                                                                                                                                                                  | TGGCTCCATC                                                                                                                                                                                                                                                                                                                                                                     |
| -86  | TCTGCAAAAG                        | GGCGCTCTGT                                                                                                         | GAGTCAGCCT<br>AP1                                                                                                                                                                               | GCTCCCCTCC                                                                                                                                                                                                                                                                                       | AGGCTTGCTC<br>p53                                                                                                                                                                                                                                                                                                                           |                                                                                                                                                                                                                                                                                                                                                                                |
| -26  | CAGCTCTCGT                        | TTCCAATGCA                                                                                                         | CGTACAGCCC                                                                                                                                                                                      | GTACACACCG                                                                                                                                                                                                                                                                                       | TGTGCTGGGA                                                                                                                                                                                                                                                                                                                                  | CACCCCACAG                                                                                                                                                                                                                                                                                                                                                                     |
|      | -446 -386 -326 -266 -206 -146 -86 | -446 GGCTCCCCTA AP2 -386 ACATGAGCTG -326 CCCTTCTGTG -266 CTGGGTGGTG -206 CCATGGCCCC -146 AGCTTTGGTA -86 TCTGCAAAAG | -446 GGCTCCCCTA GCAGCCTGCC AP2 -386 ACATGAGCTG CTTTCCCTCT -326 CCCTTCTGTG CCTGGAGCTG -266 CTGGGTGGTG CCAGGGAGAG -206 CCATGGCCCC GATAACCTTC -146 AGCTTTGGTA TGGGGGAGAG -86 TCTGCAAAAG GGCGCTCTGT | -446 GGCTCCCCTA GCAGCCTGCC CTACCTCTTT AP2 AP2  -386 ACATGAGCTG CTTTCCCTCT CAGCCAGAGG  -326 CCCTTCTGTG CCTGGAGCTG GGAAGCAGGC  -266 CTGGGTGGTG CCAGGGAGAG CCTCCATAGT  -206 CCATGGCCCC GATAACCTTC TGCCTGTGCA  -146 AGCTTTGGTA TGGGGGAGAG GGCACAGGGC  -86 TCTGCAAAAG GGCGCTCTGT GAGTCAGCCT AP1.  *** | -446 GGCTCCCCTA GCAGCCTGCC CTACCTCTTT ACCTGCTTCC AP2 CTTCCCTCT CAGCCAGAGG ACATGGGGGG -326 CCCTTCTGTG CCTGGAGCTG GGAAGCAGGC CAGGGTTAGC -266 CTGGGTGGT CCAGGGAGAG CCTGCATAGT GCCAGGTGGT -206 CCATGGCCCC GATAACCTTC TGCCTGTGCA CACACCTGCC -146 AGCTTTGGTA TGGGGGAGAG GGCACAGGGC CAGACAAACC -86 TCTGCAAAAG GGCGCTCTGT GAGTCAGCCT GCTCCCCTCC AP1 | -386 ACATGAGCTG CTTTCCCTCT CAGCCAGAGG ACATGGGGGG CCCCAGCTCC -326 CCCTTCTGTG CCTGGAGCTG GGAAGCAGGC CAGGGTTAGC TGAGGCTGGC -266 CTGGGTGGTG CCAGGGAGAG CCTGCATAGT GCCAGGTGGT GCCTTGGGTT -206 CCATGGCCCC GATAACCTTC TGCCTGTGCA CACACCTGCC CCTCACTCCA -146 AGCTTTGGTA TGGGGGAGAG GGCACAGGGC CAGACAAACC TGTGAGACTT -86 TCTGCAAAAG GGCGCTCTGT GAGTCAGCCT GCTCCCCTCC AGGCTTGCTC AP1 *** |

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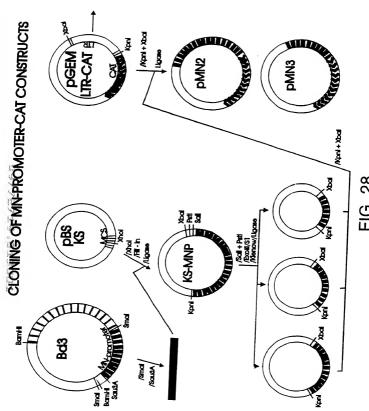


FIG. 28



